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OK protein - protein search, using sv model

Run on: August 28, 2002, 12:57:09 ; Search time 12.99 seconds  
(without alignments)  
69.573 Million cell updates/sec

Title: US-09-900-147-1  
Perfect score: 186  
Sequence: 1 KNIRRRVTDALVLMANNISSEKKEIKWIGLPTNSA 37

Scoring Table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 231628 seqs, 24425594 residues

Total number of hits satisfying chosen parameters: 231628

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Listing first 45 summaries

Database : Issued Patents AA.\*  
1: /cgn2.6/ptodata/2/1aa/5A.COMB.pap.\*  
2: /cgn2.6/ptodata/2/1aa/5B.COMB.pap.\*  
3: /cgn2.6/ptodata/2/1aa/6A.COMB.pap.\*  
4: /cgn2.6/ptodata/2/1aa/6B.COMB.pap.\*  
5: /cgn2.6/ptodata/2/1aa/PCTUS.COMB.pap.\*  
6: /cgn2.6/ptodata/2/1aa/Docfiles1.pap.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	186	100.0	37	4	US-09-308-935-1
2	186	100.0	72	2	US-08-428-131-11
3	186	100.0	72	4	US-09-078-596-11
4	186	100.0	410	2	US-08-723-415B-10
5	186	100.0	410	2	US-08-723-415B-11
6	186	100.0	410	2	US-08-428-131-2
7	186	100.0	410	2	US-08-602-846-2
8	186	100.0	410	4	US-09-078-596-2
9	186	100.0	410	4	US-09-189-627A-10
10	186	100.0	410	4	US-09-189-627A-11
11	182	97.8	369	2	US-08-723-415B-4
12	182	97.8	369	4	US-09-189-627A-4
13	182	97.8	370	2	US-08-723-415B-6
14	182	97.8	370	4	US-09-189-627A-6
15	182	97.8	385	2	US-08-723-415B-8
16	182	97.8	385	4	US-09-189-627A-8
17	182	97.8	466	2	US-08-723-415B-2
18	182	97.8	466	4	US-09-189-627A-2
19	152	81.7	30	4	US-09-308-935-6
20	101	54.3	20	4	US-09-308-935-4
21	91	48.9	19	4	US-09-308-935-3
22	83	44.6	19	4	US-09-308-935-15
23	81	43.5	17	2	US-08-428-131-13
24	81	43.0	17	4	US-09-078-596-13
25	80	43.0	15	4	US-09-308-935-10
26	79	42.5	19	4	US-09-308-935-17
27	77	41.4	19	4	US-09-308-935-16

28	76	40.9	16	4	US-09-308-935-5
29	71.5	38.4	73	2	US-08-428-131-12
30	71.5	38.4	73	4	US-09-078-596-12
31	71.5	38.4	196	2	US-08-481-814A-9
32	71.5	38.4	437	1	US-08-136-119-4
33	71.5	38.4	437	1	US-07-892-721-2
34	71.5	38.4	437	2	US-08-723-415B-13
35	71.5	38.4	437	2	US-08-481-814A-6
36	71.5	38.4	437	2	US-08-481-814A-7
37	71.5	38.4	437	3	US-08-801-092-1
38	71.5	38.4	437	4	US-09-189-627A-13
39	71.5	38.4	476	2	US-08-139-937-14
40	71.5	38.4	476	5	PCT-0593-11310-14
41	69.5	37.4	437	1	US-08-136-119-2
42	69.5	37.4	437	2	US-08-481-814A-7
43	69	37.1	14	4	US-09-308-935-11
44	65.5	35.2	413	2	US-08-481-814A-8
45	65.5	35.2	413	3	US-08-836-582-2

ALIGNMENTS

RESULT 1 *Surigial stent*

US-09-308-935-1  
; Sequence 1, Application US/09308935  
; Patent No. 6268334  
; GENERAL INFORMATION:  
; APPLICANT: La Thangue, Nicholas B  
; APPLICANT: Bandara, Laseantha R  
; TITLE OF INVENTION: Peptide antagonists of DP transcription factors  
; FILE REFERENCE: 620-67  
; CURRENT APPLICATION NUMBER: US/09/308.935  
; EARLIER FILING DATE: 1999-05-27  
; EARLIER APPLICATION NUMBER: PCT/GB97/03506  
; EARLIER FILING DATE: 1997-12-22  
; EARLIER APPLICATION NUMBER: GB 9626589.7  
; NUMBER OF SEQ ID NOS: 18  
; SOFTWARE: Patentin Ver. 2.1  
; SEQ ID NO 1  
; LENGTH: 37  
; TYPE: PPT  
; ORGANISM: Artificial Sequence  
; FEATURE:  
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic

US-09-308-935-1

Query Match 100.0% Score 186; DB 4; Length 37;  
Best Local Similarity 100.0%; Pred. No. 1.3e-21;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 1 KNIRRRVTDALVLMANNISSEKKEIKWIGLPTNSA 37  
Db 1 KNIRRRVTDALVLMANNISSEKKEIKWIGLPTNSA 37

RESULT 2 *claims on A*

US-08-428-131-11  
; Sequence 11, Application US/08428131  
; Patent No. 5863757  
; GENERAL INFORMATION:  
; APPLICANT: La Thangue, Nicholas Barrie  
; TITLE OF INVENTION: Transcription Factor DP-1  
; NUMBER OF SEQUENCES: 14  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Nixon & Vanderhye  
; STREET: 1100 No. 5863757th Glebe Road, 8th Floor  
; CITY: Arlington  
; STATE: Virginia  
; COUNTRY: U.S.A.  
; ZIP: 22201-4714

COMPUTER READABLE FORM:  
MEDIUM TYPE: floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent in Release #1.0, Version #1.25 (EPO)  
CURRENT APPLICATION DATA:  
FILING DATE: 23-JUN-1995  
CLASSIFICATION: 514  
ATTORNEY/AGENT INFORMATION:  
NAME: Arthur R. Crawford  
REGISTRATION NUMBER: 25,327  
REFERENCE/DOCKET NUMBER: 117-181  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (703) 816-4000  
TELEFAX: (703) 816-4000  
INFORMATION FOR SEQ ID NO: 11:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 72 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-428-131-11

Query Match 100.0%; Score 186; DB 2; Length 72;  
Best Local Similarity 100.0%; Pred. No. 3e-21;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 KNIRRVYDALNVLAMNIISKEKEIKWIGLPTNSA 37  
DB 4 KNIRRVYDALNVLAMNIISKEKEIKWIGLPTNSA 40

RESULT 3  
US-09-078-596-11  
Sequence 11, Application US/09078536  
Patent No. 6150116  
GENERAL INFORMATION:  
APPLICANT: La Thangue, Nicholas Barrie  
TITLE OF INVENTION: Transcription Factor DP-1  
NUMBER OF SEQUENCES: 14  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Nixon & Vanderrhye  
STREET: 1100 No. 6150116th Glebe Road, 8th Floor  
CITY: Arlington  
STATE: Virginia  
COUNTRY: U.S.A.  
ZIP: 22001-4714  
COMPUTER READABLE FORM:  
MEDIUM TYPE: floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent in Release #1.0, Version #1.25 (EPO)  
CURRENT APPLICATION DATA:  
FILING DATE: 23-JUN-1995  
CLASSIFICATION:  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US/08/428,131  
FILING DATE: 23-JUN-1995  
ATTORNEY/AGENT INFORMATION:  
NAME: Arthur R. Crawford  
REGISTRATION NUMBER: 25,327  
REFERENCE/DOCKET NUMBER: 117-181  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (703) 816-4000  
TELEFAX: (703) 816-4000  
INFORMATION FOR SEQ ID NO: 11:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 72 amino acids  
TYPE: amino acid

COMPUTER READABLE FORM:  
MEDIUM TYPE: floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent in Release #1.0, Version #1.25 (EPO)  
CURRENT APPLICATION DATA:  
FILING DATE: 23-JUN-1995  
CLASSIFICATION:  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US/08/428,131  
FILING DATE: 23-JUN-1995  
ATTORNEY/AGENT INFORMATION:  
NAME: Arthur R. Crawford  
REGISTRATION NUMBER: 25,327  
REFERENCE/DOCKET NUMBER: 117-181  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (703) 816-4000  
TELEFAX: (703) 816-4000  
INFORMATION FOR SEQ ID NO: 11:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 72 amino acids  
TYPE: amino acid

STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-09-078-596-11

Query Match 100.0%; Score 186; DB 4; Length 72;  
Best Local Similarity 100.0%; Pred. No. 3e-21;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 KNIRRVYDALNVLAMNIISKEKEIKWIGLPTNSA 37  
DB 4 KNIRRVYDALNVLAMNIISKEKEIKWIGLPTNSA 40

RESULT 4  
US-08-723-4158-10  
Sequence 10, Application US/087234158  
Patent No. 5859199  
GENERAL INFORMATION:  
APPLICANT: Lathangue, Nicholas B.  
APPLICANT: Gelaluna, Susana  
TITLE OF INVENTION: TRANSCRIPTION FACTOR DP-3 AND ISOFORMS  
TITLE OF INVENTION: THEREOF  
NUMBER OF SEQUENCES: 21  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: NIXON & VANDERRHYE P.C.  
STREET: 1100 No. 5859199th Glebe Rd. 8th floor  
CITY: Arlington  
STATE: VA  
COUNTRY: USA  
ZIP: 22201-4741  
COMPUTER READABLE FORM:  
MEDIUM TYPE: floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent in Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/723,4158  
FILING DATE: 30-SEP-1996  
CLASSIFICATION: 435  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: GB 9610195.1  
FILING DATE: 15-MAY-1996  
ATTORNEY/AGENT INFORMATION:  
NAME: Crawford, Arthur R.  
REGISTRATION NUMBER: 25,327  
REFERENCE/DOCKET NUMBER: 117-220  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 703-816-4000  
TELEFAX: 703-816-4100  
INFORMATION FOR SEQ ID NO: 10:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 410 amino acids  
TYPE: amino acid  
STRANDEDNESS:  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-723-4158-10

Query Match 100.0%; Score 186; DB 2; Length 410;  
Best Local Similarity 100.0%; Pred. No. 2.3e-20;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 KNIRRVYDALNVLAMNIISKEKEIKWIGLPTNSA 37  
DB 163 KNIRRVYDALNVLAMNIISKEKEIKWIGLPTNSA 199

RESULT 5  
US-08-723-4158-11  
Sequence 11, Application US/087234158

Patent No. 5859199  
GENERAL INFORMATION:  
APPLICANT: LaThangue, Nicholas B.  
APPLICANT: deLaluna, Susana  
TITLE OF INVENTION: TRANSCRIPTION FACTOR DP-3 AND ISOFORMS  
TITLE OF INVENTION: THEREOF  
NUMBER OF SEQUENCES: 21  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: NIXON & VANDERHYE P.C.  
STREET: 1100 No. 5859199th Glebe Rd. 8th floor  
CITY: Arlington  
STATE: VA  
COUNTRY: USA  
ZIP: 22201-4714  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent In Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/723,415B  
FILING DATE: 30-SEP-1996  
CLASSIFICATION: 435  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: GB 9610195.1  
FILING DATE: 15-MAY-1996  
ATTORNEY/AGENT INFORMATION:  
NAME: Crawford, Arthur R.  
REGISTRATION NUMBER: 25,327  
REFERENCE/DOCKET NUMBER: 117-220  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 703-816-4000  
TELEFAX: 703-816-4100  
INFORMATION FOR SEQ ID NO: 11:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 410 amino acids  
TYPE: amino acid  
STRANDEDNESS:  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-723-415B-11

Query Match 100.0%; Score 186; DB 2; Length 410;  
Best Local Similarity 100.0%; Pred. No. 2.3e-20;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 1 KNIRRVYDALVLMANNIISKEKKEIKWIGLPTNSA 37  
Db 163 KNIRRVYDALVLMANNIISKEKKEIKWIGLPTNSA 199

RESULT 6  
US-08-428-131-2  
Sequence 2, Application US/08/428131  
Patent No. 5863717  
GENERAL INFORMATION:  
APPLICANT: LaThangue, Nicholas Barrie  
TITLE OF INVENTION: TRANSCRIPTION FACTOR DP-1  
NUMBER OF SEQUENCES: 14  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: NIXON & VANDERHYE  
STREET: 1100 No. 5863757th Glebe Road, 8th floor  
CITY: Arlington  
STATE: Virginia  
COUNTRY: U.S.A.  
ZIP: 22201-4714  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent In Release #1.0, Version #1.25 (EPO)  
CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/428,131  
FILING DATE: 23-JUN-1995  
CLASSIFICATION: 514  
ATTORNEY/AGENT INFORMATION:  
NAME: Arthur R. Crawford  
REGISTRATION NUMBER: 25,327  
REFERENCE/DOCKET NUMBER: 117-181  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (703) 816-4000  
TELEFAX: (703) 816-4100  
INFORMATION FOR SEQ ID NO: 2:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 410 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-428-131-2

Query Match 100.0%; Score 186; DB 2; Length 410;  
Best Local Similarity 100.0%; Pred. No. 2.3e-20;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 1 KNIRRVYDALVLMANNIISKEKKEIKWIGLPTNSA 37  
Db 163 KNIRRVYDALVLMANNIISKEKKEIKWIGLPTNSA 199

RESULT 7  
US-08-602-846-2  
Sequence 2, Application US/08/602846  
Patent No. 5871901  
GENERAL INFORMATION:  
APPLICANT: LaThangue, Nicholas B  
TITLE OF INVENTION: ASSAY FOR INHIBITORS OF DP-1 AND OTHER DP  
TITLE OF INVENTION: PROTEINS.  
NUMBER OF SEQUENCES: 3  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Nixon & Vanderhye PC  
STREET: 8th floor, 1100 No. 5871901th Glebe Road  
CITY: Arlington  
STATE: Virginia  
COUNTRY: USA  
ZIP: 22201-4714  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent In Release #1.0  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/602,846  
FILING DATE: 26-FEB-1996  
CLASSIFICATION: 435  
ATTORNEY/AGENT INFORMATION:  
NAME: CRAWFORD, ARTHUR R.  
REGISTRATION NUMBER: 25,327  
REFERENCE/DOCKET NUMBER: 620-12  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (703) 816-4000  
TELEFAX: (703) 816-4100  
INFORMATION FOR SEQ ID NO: 2:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 410 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-602-846-2

Query Match 100.0%; Score 186; DB 2; Length 410;  
Best Local Similarity 100.0%; Pred. No. 2.3e-20;  
Matches 37; Conservative 0; Mismatches 0; Indels 0; Gaps 0;



APPLICATION NUMBER: US/08/723,415B  
FILING DATE: 30-SEP-1996  
CLASSIFICATION: 435  
PRIOR APPLICATION DATA:  
PRIOR APPLICATION NUMBER: GB 9610195.1  
FILING DATE: 15-MAY-1996  
ATTORNEY/AGENT INFORMATION:  
NAME: Crawford, Arthur R.  
REGISTRATION NUMBER: 25,327  
REFERENCE/DOCKET NUMBER: 117-220  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 703-816-4000  
TELEFAX: 703-816-4100  
INFORMATION FOR SEQ ID NO: 4:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 369 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-723-415B-4

Query Match 97.88; Score 182; DB 2; Length 369;  
Best Local Similarity 97.38; Pred. No. 8.4e-20;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

OY 1 KNIRRVYDALVLMNNIISKKEIKWIGLPTNSA 37  
:|||||  
DB 101 ENIRRVYDALVLMNNIISKKEIKWIGLPTNSA 137

## RESULT 12

US-09-189-627A-4  
Sequence 4, Application US/09189627A  
Patent No. 6159691  
GENERAL INFORMATION:  
APPLICANT: La Thangue, Nicholas  
APPLICANT: de la Luna, Susana  
TITLE OF INVENTION: TRANSCRIPTION FACTOR DP-3 AND ISOFORMS THEREOF  
FILE REFERENCE: 620-54  
CURRENT APPLICATION NUMBER: US/09/189,627A  
PRIOR FILING DATE: 1998-11-10  
PRIOR APPLICATION NUMBER: 08/723,415  
PRIOR FILING DATE: 1996-09-30  
PRIOR APPLICATION NUMBER: GB 9610195  
PRIOR FILING DATE: 1996-05-15  
NUMBER OF SEQ ID NOS: 25  
SOFTWARE: PatentIn Ver. 2.0  
SEQ ID NO 4  
LENGTH: 369  
TYPE: PRT  
ORGANISM: mouse  
US-09-189-627A-4

Query Match 97.88; Score 182; DB 4; Length 369;  
Best Local Similarity 97.38; Pred. No. 8.4e-20;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

OY 1 KNIRRVYDALVLMNNIISKKEIKWIGLPTNSA 37  
:|||||  
DB 101 ENIRRVYDALVLMNNIISKKEIKWIGLPTNSA 137

## RESULT 13

US-08-723-415B-6  
Sequence 6, Application US/08723415B  
Patent No. 5839199  
GENERAL INFORMATION:  
APPLICANT: La Thangue, Nicholas B.  
APPLICANT: de la Luna, Susana  
TITLE OF INVENTION: TRANSCRIPTION FACTOR DP-3 AND ISOFORMS THEREOF

NUMBER OF SEQUENCES: 21  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: NIXON & VANDERHYTE P.C.  
STREET: 1100 NO. 5839199th Giede Rd. 8th floor  
CITY: Arlington  
STATE: VA  
COUNTRY: USA  
ZIP: 22201-4741  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/723,415B  
FILING DATE: 30-SEP-1996  
CLASSIFICATION: 435  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: GB 9610195.1  
FILING DATE: 15-MAY-1996  
ATTORNEY/AGENT INFORMATION:  
NAME: Crawford, Arthur R.  
REGISTRATION NUMBER: 25,327  
REFERENCE/DOCKET NUMBER: 117-220  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 703-816-4000  
TELEFAX: 703-816-4100  
INFORMATION FOR SEQ ID NO: 6:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 370 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-723-415B-6

Query Match 97.88; Score 182; DB 2; Length 370;  
Best Local Similarity 97.38; Pred. No. 8.4e-20;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

OY 1 KNIRRVYDALVLMNNIISKKEIKWIGLPTNSA 37  
:|||||  
DB 102 ENIRRVYDALVLMNNIISKKEIKWIGLPTNSA 138

## RESULT 14

US-09-189-627A-6  
Sequence 6, Application US/09189627A  
Patent No. 6159691  
GENERAL INFORMATION:  
APPLICANT: La Thangue, Nicholas  
APPLICANT: de la Luna, Susana  
TITLE OF INVENTION: TRANSCRIPTION FACTOR DP-3 AND ISOFORMS THEREOF  
FILE REFERENCE: 620-54  
CURRENT APPLICATION NUMBER: US/09/189,627A  
PRIOR FILING DATE: 1998-11-10  
PRIOR APPLICATION NUMBER: 08/723,415  
PRIOR FILING DATE: 1996-09-30  
PRIOR APPLICATION NUMBER: GB 9610195  
PRIOR FILING DATE: 1996-05-15  
NUMBER OF SEQ ID NOS: 25  
SOFTWARE: PatentIn Ver. 2.0  
SEQ ID NO 6  
LENGTH: 370  
TYPE: PRT  
ORGANISM: mouse  
US-09-189-627A-6

Query Match 97.88; Score 182; DB 4; Length 370;  
Best Local Similarity 97.38; Pred. No. 8.4e-20;  
Matches 36; Conservative 1; Mismatches 0; Indels 0; Gaps 0;